

# SEQUENCE LISTING

<110> Novozymes Biopharma UK Limited  
 Sleep, Darrell  
  
 <120> Gene and Polypeptide Sequences  
  
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 <141> 2005-07-08  
  
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 <151> 2002-07-23  
  
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Tyr Ser															

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Met	Lys	Trp	Val	Xaa	Xaa	Xaa	Xaa	Xaa	Leu	Phe	Leu	Phe	Ser	Ser	Ala
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ccattcgaag	atcacgtcaa	gttgggtcaac	gaagttaccg	aattcgctaa	gacttggtgt	180
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gaaccagaaa	gaaacgaatg	tttcttgcaa	cacaaggacg	acaacccaaa	cttgccaaga	360
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aagcaaagat	tgaagtgtgc	ttccttgcaa	aagttcgggt	aaagagcttt	caaggcttgg	660
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gaagatcacg	tcaagttggg	caacgaagtt	accgaattcg	ctaagacttg	tgttgctgac	240
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tgtactgaat	ctttgggttaa	cagaagacca	tgtttctctg	ctttggaagt	cgacgaaact	1560
tacgttccaa	aggaattcaa	cgctgaaact	ttcaccttcc	acgctgatat	ctgtaccttg	1620
tccgaaaagg	aaagacaaat	taagaagcaa	actgctttgg	ttgaattggg	caagcacaa	1680
ccaaaggcta	ctaaggaaca	attgaaggct	gtcatggatg	atttcgctgc	tttcggtgaa	1740
aagtgttgta	aggctgatga	taaggaaact	tgtttcgctg	aagaaggtaa	gaagttggtc	1800
gctgcttccc	aagctgcttt	gggttttg				1827

<210> 22

<211> 1827

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide leader sequence and the mature human albumin coding region

<400> 22

atgaagtggg	taagctttat	ttcccttctt	tttctcttta	gctcggttta	ttccaggagc	60
ttggataaaa	gagatgcaca	caagagttag	gttgctcatc	ggtttaaaga	tttgggagaa	120
gaaaatttca	aagccttggg	gttgattgcc	tttgcctcag	atcttcagca	gtgtccattt	180
gaagatcatg	taaaattagt	gaatgaagta	actgaatttg	caaaaacatg	tgttgctgat	240
gagtcagctg	aaaattgtga	caaatcactt	catacccttt	ttggagacaa	attatgcaca	300
gttgcaactc	ttcgtgaaac	ctatggtgaa	atggctgact	gctgtgcaaa	acaagaacct	360
gagagaaatg	aatgcttctt	gcaacacaaa	gatgacaacc	caaacctccc	ccgattgggtg	420
agaccagagg	ttgatgtgat	gtgcactgct	tttcatgaca	atgaagagac	atttttgaaa	480
aaatacttat	atgaaattgc	cagaagacat	ccttactttt	atgccccgga	actccttttc	540
tttgctaaaa	ggtataaagc	tgcttttaca	gaatgttgcc	aagctgctga	taaagctgcc	600
tgctgttgct	caaagctcga	tgaacttcgg	gatgaaggga	aggcttcgtc	tgccaaacag	660
agactcaagt	gtgccagtct	ccaaaaattt	ggagaaagag	ctttcaaagc	atgggcagta	720
gctcgcttga	gccagagatt	tcccaaagct	gagtttgag	aagtttccaa	gttagtgaca	780
gatcttacca	aagtccacac	ggaatgctgc	catggagatc	tgcttgaatg	tgctgatgac	840
agggcgagcc	ttgccaagta	tatctgtgaa	aatcaagatt	cgatctccag	taaactgaag	900
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gagatgcctg	ctgacttgcc	ttcattagct	gctgattttg	ttgaaagtaa	ggatgtttgc	1020
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cagcttggag	agtacaaatt	ccagaatgcg	ctattagtgc	gttacaccaa	gaaagtaccc	1320
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ccaaggcaa	caaaagagca	actgaaagct	gttatggatg	atttcgcagc	ttttgtagag	1740
aagtgtgca	aggctgacga	taaggagacc	tgcttttgccg	aggagggtaa	aaaacttggt	1800
gctgcaagtc	aagctgcctt	aggctta				1827

<210> 23  
 <211> 47  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide leader sequence

<400> 23		
ctaaagagaa	aaagaatgga gacgatgaat acccacttca tctttgc	47

<210> 24  
 <211> 72  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide leader sequence

<400> 24		
atgaagtggg	tattcatcgt ctccattctt tttctcttta gctcggctta ttccaggagc	60
ttggataaaa	ga	72

<210> 25  
 <211> 1827  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide leader sequence and mature human albumin coding region

<400> 25		
atgaagtggg	tattcatcgt ctccattctt tttctcttta gctcggctta ttccaggagc	60
ttggataaaa	gagatgcaca caagagtggg gttgctcatc ggttttaaaga tttgggagaa	120
gaaaatttca	aagccttggt gttgattgcc tttgctcagt atcttcagca gtgtccattt	180
gaagatcatg	taaaattagt gaatgaagta actgaatttg caaaaacatg tggtgctgat	240
gagtcagctg	aaaattgtga caaatcactt catacccttt ttggagacaa attatgcaca	300
gttgcaactc	ttcgtgaaac ctatggtgaa atggctgact gctgtgcaaa acaagaacct	360

gagagaaatg	aatgcttctt	gcaacacaaa	gatgacaacc	caaacctccc	ccgattgggtg	420
agaccagagg	ttgatgtgat	gtgcactgct	tttcatgaca	atgaagagac	atTTTTgaaa	480
aaatacttat	atgaaattgc	cagaagacat	ccttactttt	atgccccgga	actccttttc	540
tttgctaaaa	ggtataaaagc	tgctttttaca	gaatgttgcc	aagctgctga	taaagctgcc	600
tgctgtttgc	caaagctcga	tgaacttcgg	gatgaaggga	aggcttcgtc	tgccaaacag	660
agactcaagt	gtgccagtct	ccaaaaatTT	ggagaaaagag	ctttcaaagc	atgggagcagta	720
gctcgctga	gccagagatt	tcccaaagct	gagtttgag	aagtttccaa	gttagtgaca	780
gatcttacca	aagtccacac	ggaatgctgc	catggagatc	tgcttgaatg	tgctgatgac	840
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gaatgctgtg	aaaaacctct	gttggaaaaa	tcccaactgca	ttgccgaagt	ggaaaatgat	960
gagatgcctg	ctgacttgcc	ttcattagct	gctgattttg	ttgaaagtaa	ggatgtttgc	1020
aaaaactatg	ctgaggcaaa	ggatgtcttc	ctgggcatgt	ttttgtatga	atatgcaaga	1080
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cagcttgagg	agtacaaatt	ccagaatgcg	ctattagttc	gttacaccaa	gaaagtaccc	1320
caagtgtcaa	ctccaactct	tgtagaggtc	tcaagaaacc	taggaaaagt	gggcagcaaa	1380
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tctgagaagg	agagacaaat	caagaaacaa	actgcacttg	ttgagctcgt	gaaacacaag	1680
cccaaggcaa	caaaagagca	actgaaagct	gttatggatg	atttcgcagc	ttttgtagag	1740
aagtgtgca	aggctgacga	taaggagacc	tgctttgccg	aggagggtaa	aaaacttggt	1800
gctgcaagtc	aagctgcctt	aggctta				1827

<210> 26  
 <211> 1827  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide plasmid sequence

<400>	26					
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gaaaacttca	aggctttggt	cttgatcgct	ttcgctcaat	acttgcaaca	atgtccattc	180
gaagatcacg	tcaagttggt	caacgaagtt	accgaattcg	ctaagacttg	tggtgctgac	240
gaatctgctg	aaaactgtga	caagtccttg	cacaccttgt	tcggtgataa	gttgtgtact	300
gttgctacct	tgagagaaac	ctacggtgaa	atggctgact	gttgtgctaa	gcaagaacca	360
gaaagaaacg	aatgtttctt	gcaacacaag	gacgacaacc	caaacttgcc	aagattgggt	420
agaccagaag	ttgacgtcat	gtgtactgct	ttccacgaca	acgaagaaac	cttcttgaag	480
aagtacttgt	acgaaattgc	tagaagacac	ccatacttct	acgctccaga	attgtttgtc	540
ttcgctaaga	gatacaaggc	tgctttcacc	gaatgttgct	aagctgctga	taaggctgct	600
tgtttggttg	caaagttgga	tgaattgaga	gacgaaggta	aggcttcttc	cgctaagcaa	660
agattgaagt	gtgcttcctt	gcaaaagttc	ggtgaaagag	ctttcaaggc	ttgggctgtc	720
gctagattgt	ctcaaagatt	cccaaaggct	gaattcgctg	aagtttctaa	gttggttact	780
gacttgacta	aggttcacac	tgaatgttgt	cacggtgact	tggtggaatg	tgctgatgac	840
agagctgact	tggttaagta	catctgtgaa	aaccaagact	ctatctcttc	caagttgaag	900
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gaaatgccag	ctgacttgcc	atctttggct	gctgacttcg	ttgaatctaa	ggacgtttgt	1020
aagaactacg	ctgaagctaa	ggacgtcttc	ttgggtatgt	tcttgtagca	atacgctaga	1080
agacaccacg	actactccgt	tgtcttggtg	ttgagattgg	ctaagacct	cgaaactacc	1140
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caattgggtg	aatacaagtt	ccaaaacgct	ttgttggtta	gatacactaa	gaagggtccca	1320

caagtctcca	ccccaaacttt	ggttgaagtc	tctagaaact	tgggtaaggt	cggttctaag	1380
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aagtgttgta	aggctgatga	taaggaaact	tgtttcgctg	aagaaggtaa	gaagttgggc	1800
gctgcttccc	aagctgcttt	gggtttg				1827

<210> 27  
 <211> 72  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide leader sequence

<400> 27	
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ttggataaga	ga
	60
	72

<210> 28  
 <211> 18  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Synthetic oligonucleotide secretion pre-sequence

<400> 28

Met	Lys	Trp	Val	Phe	Ile	Val	Ser	Ile	Leu	Phe	Leu	Phe	Ser	Ser	Ala
1				5				10					15		
Tyr	Ser														

<210> 29  
 <211> 5  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Synthetic polypeptide leader sequence

<220>  
 <221> MISC\_FEATURE  
 <222> (1)..(1)  
 <223> CAN BE EITHER Phe OR Trp OR Tyr

<220>  
 <221> MISC\_FEATURE  
 <222> (2)..(2)  
 <223> CAN BE EITHER Ile OR Leu OR Val OR Ala OR Met

<220>  
<221> MISC\_FEATURE  
<222> (3)..(3)  
<223> CAN BE EITHER Leu OR Val OR Ala OR Met

<220>  
<221> MISC\_FEATURE  
<222> (5)..(5)  
<223> CAN BE EITHER Ile OR Val OR Ala OR Met

<400> 29

Xaa Xaa Xaa Thr Xaa  
1 5

<210> 30  
<211> 15  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic polypeptide secretion pre-sequence

<400> 30

Leu Phe Leu Phe Ser Ser Ala Tyr Ser Arg Gly Val Phe Arg Arg  
1 5 10 15

<210> 31  
<211> 24  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic polypeptide secretion pre-sequence

<220>  
<221> MISC\_FEATURE  
<222> (5)..(5)  
<223> any amino acid PREFERABLY Phe

<220>  
<221> MISC\_FEATURE  
<222> (6)..(6)  
<223> any amino acid PREFERABLY Ile

<220>  
<221> MISC\_FEATURE  
<222> (7)..(7)  
<223> any amino acid PREFERABLY Val

<220>  
<221> MISC\_FEATURE  
<222> (8)..(8)

<223> any amino acid PREFERABLY Ser OR Thr

<220>

<221> MISC\_FEATURE

<222> (9)..(9)

<223> any amino acid PREFERABLY Ile

<400> 31

Met	Lys	Trp	Val	Xaa	Xaa	Xaa	Xaa	Xaa	Leu	Phe	Leu	Phe	Ser	Ser	Ala
1				5					10					15	
Tyr	Ser	Arg	Gly	Val	Phe	Arg	Arg								
			20												

<210> 32

<211> 24

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic polypeptide secretion pre-pro sequence

<400> 32

Met	Lys	Trp	Val	Phe	Ile	Val	Ser	Ile	Leu	Phe	Leu	Phe	Ser	Ser	Ala
1				5					10					15	
Tyr	Ser	Arg	Ser	Leu	Asp	Lys	Arg								
			20												

<210> 33

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic polypeptide secretion pre-sequence

<220>

<221> MISC\_FEATURE

<222> (2)..(2)

<223> CAN BE EITHER Lys OR Arg OR His

<220>

<221> MISC\_FEATURE

<222> (3)..(3)

<223> CAN BE EITHER Phe OR Trp OR Tyr

<220>

<221> MISC\_FEATURE

<222> (4)..(4)

<223> CAN BE EITHER Ile OR Leu OR Val OR Ala OR Met

<400> 33

Met Xaa Xaa Xaa

1

<210> 34  
<211> 15  
<212> DNA  
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<220>  
<223> Synthetic oligonucleotide leader sequence

<400> 34  
ttcatcgtct ccatt 15

<210> 35  
<211> 36  
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<220>  
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<400> 35  
gcatgcgggcc gcccgtaatg cggatcgtg aaagcg 36

<210> 36  
<211> 35  
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<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide primer

<400> 36  
gcataagctt acccacttca tctttgcttg tttag 35

<210> 37  
<211> 11  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide linker

<400> 37  
ttaggcttat a 11

<210> 38  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide linker

<400> 38  
ccgaatattc ga 12

<210> 39  
<211> 40  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide primer

<400> 39  
gttagaatta ggттаagctt gtttttttat tggcgatgaa 40

<210> 40  
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<220>  
<223> s. cerevisiae 5'UTR and synthetic oligonucleotide leader sequence  
and mature human albumin CDS

<400> 40

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cttgttctcc	tctgcttact	ctagatcttt	ggataagaga	gacgctcaca	agtcgcgaagt	120
cgctcacaga	ttcaaggact	tgggtgaaga	aaacttcaag	gcttttggct	tgatcgcttt	180
cgctcaatac	ttgcaacaat	gtccattcga	agatcacgct	aagttgggtca	acgaagttac	240
cgaattcgct	aagacttggt	ttgctgacga	atctgctgaa	aactgtgaca	agtccttgca	300
caccttggtc	ggtgataagt	tgtgtactgt	tgctaccttg	agagaaacct	acggtgaaat	360
ggctgactgt	tgtgctaagc	aagaaccaga	aagaaacgaa	tgtttcttgc	aacacaagga	420
cgacaaccca	aacttgccaa	gattggttag	accagaagtt	gacgtcatgt	gtactgcttt	480
ccacgacaac	gaagaaacct	tcttgaagaa	gtacttgtac	gaaattgcta	gaagacaccc	540
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atgttggtcaa	gctgctgata	aggctgcttg	tttgttgcca	aagttggatg	aattgagaga	660
cgaaggtaag	gcttcttccg	ctaagcaaag	attgaagtgt	gcttccttgc	aaaagttcgg	720
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ccaagactct	atctcttcca	agttgaagga	atgttggtgaa	aagccattgt	tggaaaagtc	960
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tttcgctgaa gaaggtaaga agttggtcgc tgcttcccaa gctgctttgg gtttgtaata	1860
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